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FIGURE 1

hRANTES	ASPYSSDT TPCCFAYIARPLPRAHIKEYFYTS GK	CSPNAVVFVTRKGRQV	CANPEKKWVREYINSLEHS
mMIP-1 α	APYGADTPTACCFSY SRKIPRQFIVDYFETSSL	CSQPGVIFLTERNRQI	CADSKETWQVEYITDLELNA
mMIP-1 β	APMGSDPFTSCCFSYTSRQLHRSFVNDYYETSSL	CSKPAVVFLTKRGRQI	CANPSEPWVTEYMSDLELN
MCP-1	LAQPDAINAPVTCYNTNRKISVQRLASVRRITSSK	CPKEAVIFKTIIVAKEI	CADPKQKVQDSMDHLDKQTQTPKT
MCP-3	SPQGLAQPVGINTSFTCCYRHFINKKIPKQRLSEYRRITSSH	CPREAVIFKTKLDKEI	CADFTQKVQDFMKHLDDKKTQTQPKL
CK β -1	TKTESSSRGPHPESECCFTYTYKI PRQRIMDYETNSQ	CSKPGIVFITXRGHSV	CTNPSDKWVQDYIKDMK
CK β -4	ASNFDCCLGYTDRILLHPKFIVGTRQLANBGCDINAIIFHTKKLSV	CSKPGVIFLTKGRRF	CANPKQTVWVYIVRLLSKKVKNN
CK β -6	VVIPSPCCMFFVSKRIPENRNVVSQLSRST	CLKGGVIFTTKGQQF	CGDEKQEWQRVMKNLDAKQKKA
CK β -7	AQVGYNKELCCLVYTSWQIFQKFI VDYSETSPQ	CPKPGVILLTKRGRQI	CADENKKRWVQK
CK β -8	ENPVLLDRFHATSADCCISYTPRSIPCSLLSESYFETNSE	CSKPGVIFLTKGRRF	CANPSDKQVQVCMENLKLDTRIKTRKN
CK β -9	SDAGGAQDCCLKYSQRKI PAKVVRSYRKQEPESLGC SI	PAIILFLPRKRSQAEEL	CADPKELKVQQLMQHLCKTPSPQKPAQ
CK β -10	ENPQGLAQPDALNVPSTCCPTFESSKKISLQRLKSYVITTSR	CPQKAVIFRTKLGKEI	CADPEKENVQNYMKHLGRKAHTLKT
CK β -11	PAPTLSGTND AEDCCLSVTQKPIPGYIVNFHYLL LKDGCRVP	PAVVFTTLRGRQL	CAPPDQPNVERIIQRLQRTSAKGRRSS
CK β -12	RSQPKVPEWVNTPTSTCCLKYTEKVLPRRLVVGTRKALN	CHLPALIIFVTRNREV	CTNPNDDWVQVEYIKDNLPLLPTRNLST
CK β -13	PYGANNMEDSVTCRDYVRYRLPLRVVVKHFYWTSDS	CPRPGVVLLTFRDKBI	CADPRVPVWVCHILNKLSSQ
NAP-1/IL-8	SAKELRCQCIKITYSKPFPHPKFIKELRVIESGPHCANTEIIVKLSDGREL	CLDDPKENWVQRVVEKFLKRAENS	
NAP-2	AELRCMCIKITYSGIHPKNIQSLEVVIGKGTGHCNQVEVLATLRDGRKI	CLDDPADPRIKCIIVQKKLAGDESAD	
hPF4	EAEEDGDLQCLCVKTTTSQVRPHITSLSEVIKAGPH	CPTAQLLATLRKGRKI	CLDLQAPLYKKILKKLES
CK α -1	VLEVYTTSLRCRCVQESSVFIPRRFIDRIQILPRGNGCCPRKEIIVMKNNK	SIVCVDPQAEWIQRMMEVLEKKR	